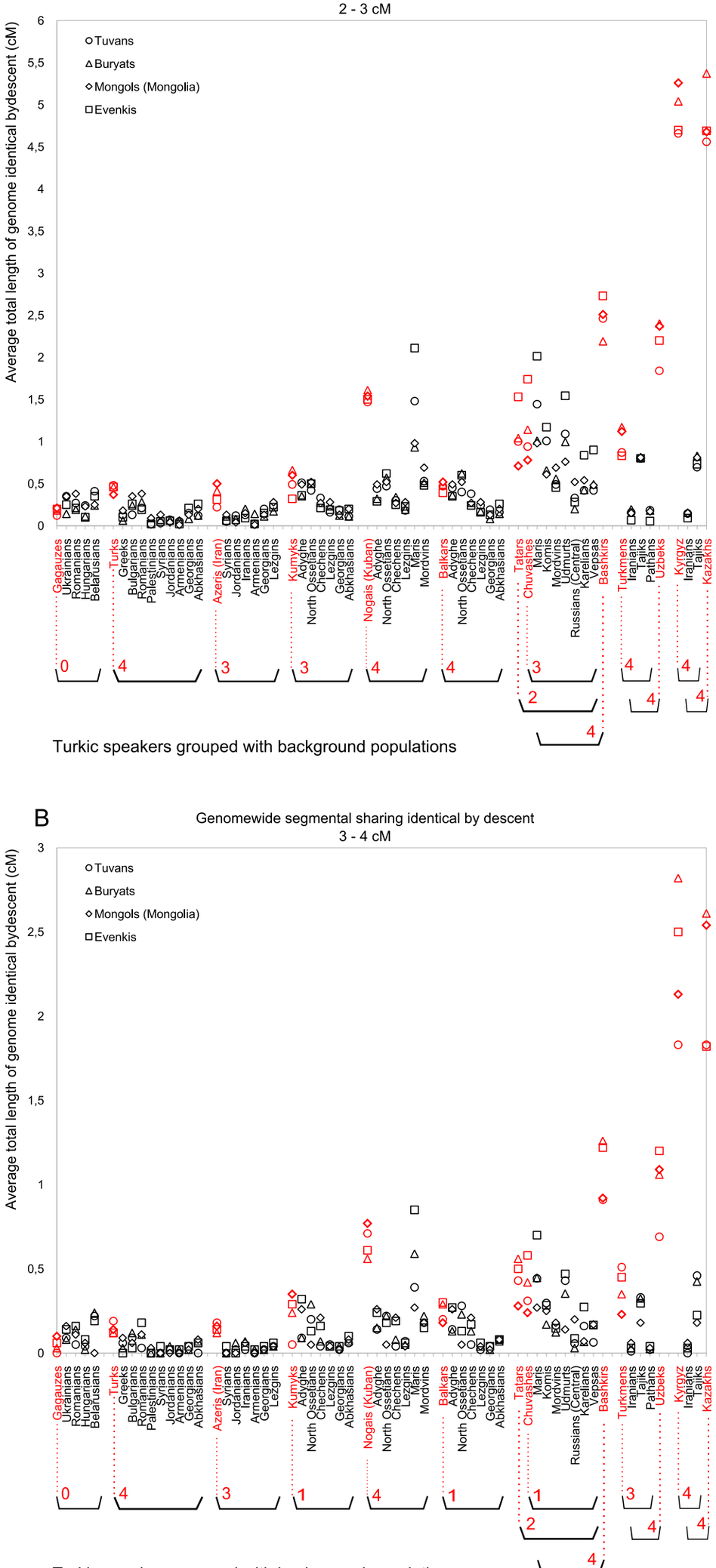
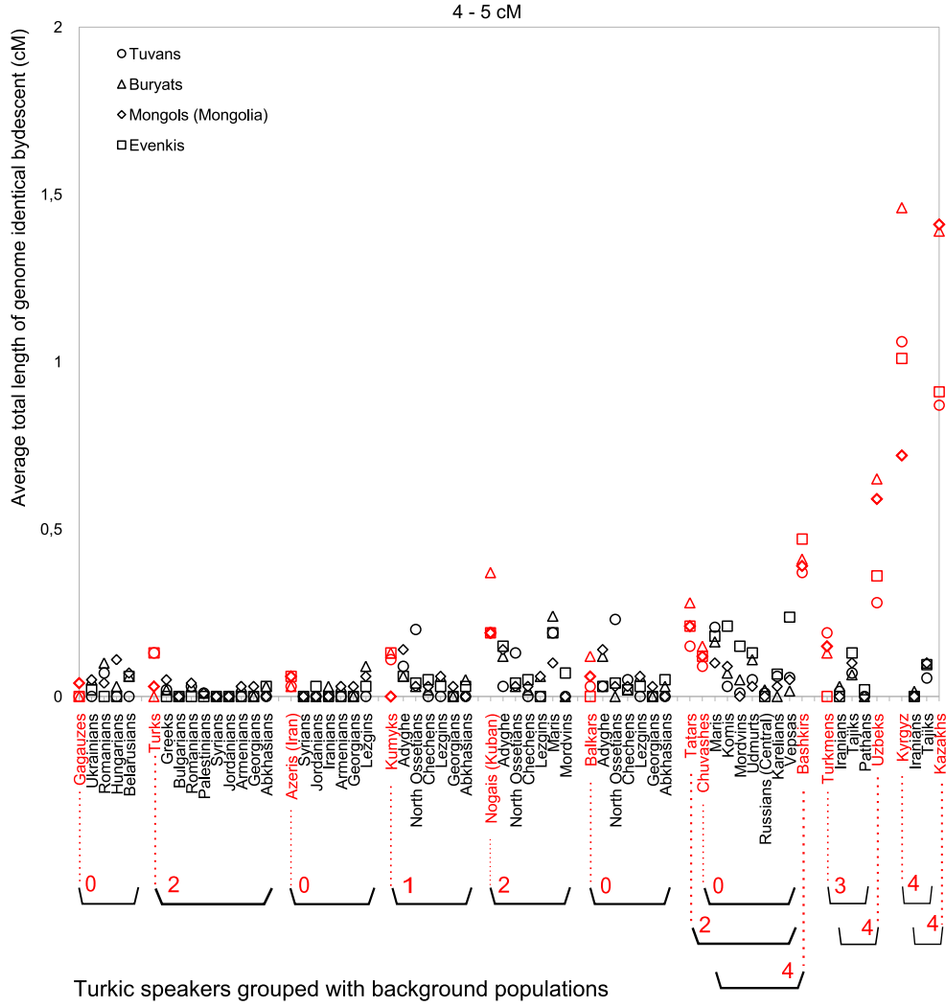
A


Genomewide segmental sharing identical by descent
2 - 3 cM
Average total length of genome identical bydescent (cM)
Tuvans
Buryats
Mongols (Mongolia)
Evenkis
Gagauzes Ukrainians Romanians Hungarians Belarusians
Turks Greeks Bulgarians Romanians Palestinians Syrians Jordanians Armenians Georgians Abkhasians
Azeris (Iran) Syrians Jordanians Iranians Armenians Georgians Lezgins
Kumyks Adyghe North Ossetians Chechens Lezgins Georgians Abkhasians
Nogais (Kuban) Adyghe North Ossetians Chechens Lezgins Maris Mordvins
Balkars Adyghe North Ossetians Chechens Lezgins Georgians Abkhasians
Tatars Chuvashes Maris Komis Mordvins Udmurts Russians (Central) Karelians Vepsas Bashkirs
Turkmens Iranians Tajiks Pathans Uzbeks
Kyrgyz Iranians Tajiks Kazakhs
Turkic speakers grouped with background populations

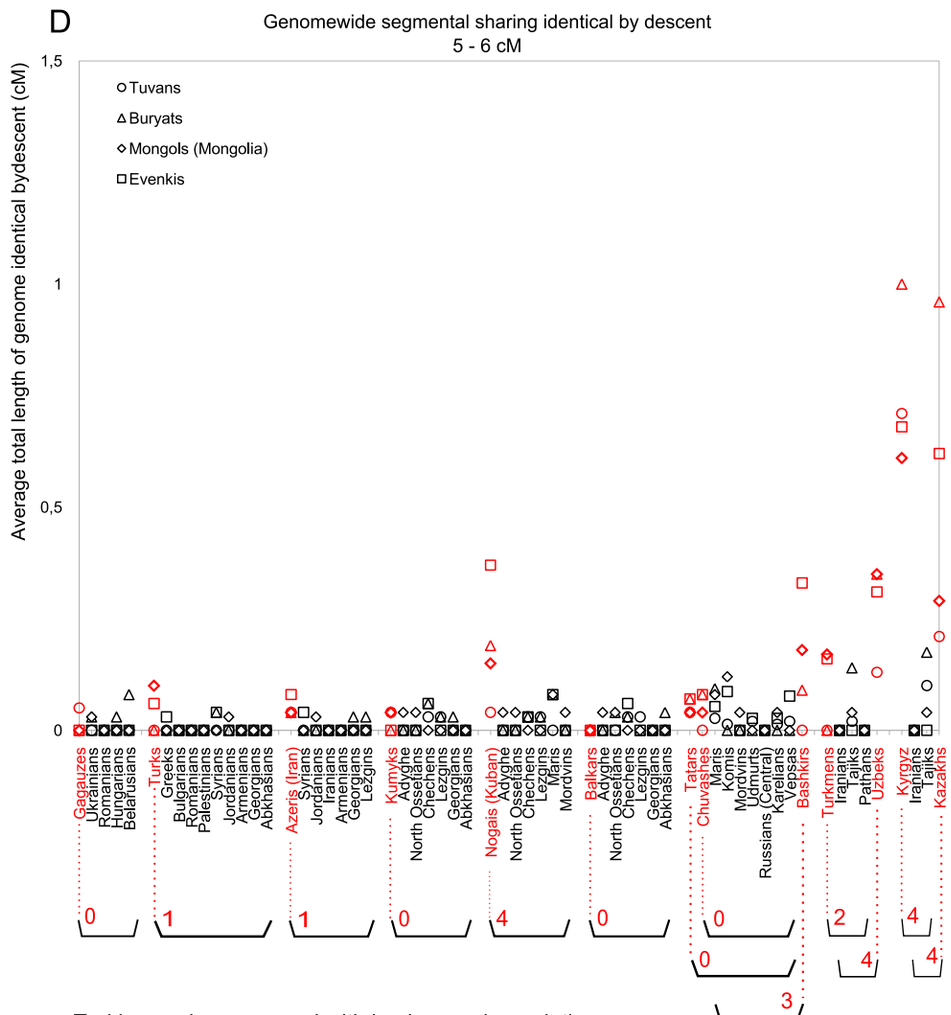
B

Genomewide segmental sharing identical by descent
3 - 4 cM
Average total length of genome identical bydescent (cM)
Tuvans
Buryats
Mongols (Mongolia)
Evenkis
Gagauzes Ukrainians Romanians Hungarians Belarusians
Turks Greeks Bulgarians Romanians Palestinians Syrians Jordanians Armenians Georgians Lezgins
Azeris (Iran) Syrians Jordanians Iranians Armenians Georgians Lezgins
Kumyks Adyghe North Ossetians Chechens Lezgins Georgians Abkhasians
Nogais (Kuban) Adyghe North Ossetians Chechens Lezgins Maris Mordvins
Balkars Adyghe North Ossetians Chechens Lezgins Georgians Abkhasians
Tatars Chuvashes Maris Komis Mordvins Udmurts Russians (Central) Karelians Vepsas Bashkirs
Turkmens Iranians Tajiks Pathans Uzbeks
Kyrgyz Iranians Tajiks Kazakhs
Turkic speakers grouped with background populations

C


Genomewide segmental sharing identical by descent
4 - 5 cM
Average total length of genome identical bydescent (cM)
Tuvans
Buryats
Mongols (Mongolia)
Evenkis
Gagauzes Ukrainians Romanians Hungarians Belarusians
Turks Greeks Bulgarians Romanians Palestinians Syrians Jordanians Armenians Georgians Lezgins
Azeris (Iran) Syrians Jordanians Iranians Armenians Georgians Lezgins
Kumyks Adyghe North Ossetians Chechens Lezgins Georgians Abkhasians
Nogais (Kuban) Adyghe North Ossetians Chechens Lezgins Maris Mordvins
Balkars Adyghe North Ossetians Chechens Lezgins Georgians Abkhasians
Tatars Chuvashes Maris Komis Mordvins Udmurts Russians (Central) Karelians Vepsas Bashkirs
Turkmens Iranians Tajiks Pathans Uzbeks
Kyrgyz Iranians Tajiks Kazakhs
Turkic speakers grouped with background populations

D


Genomewide segmental sharing identical by descent
5 - 6 cM
Average total length of genome identical bydescent (cM)
Tuvans
Buryats
Mongols (Mongolia)
Evenkis
Gagauzes Ukrainians Romanians Hungarians Belarusians
Turks Greeks Bulgarians Romanians Palestinians Syrians Jordanians Armenians Georgians Lezgins
Azeris (Iran) Syrians Jordanians Iranians Armenians Georgians Lezgins
Kumyks Adyghe North Ossetians Chechens Lezgins Georgians Abkhasians
Nogais (Kuban) Adyghe North Ossetians Chechens Lezgins Maris Mordvins
Balkars Adyghe North Ossetians Chechens Lezgins Georgians Abkhasians
Tatars Chuvashes Maris Komis Mordvins Udmurts Russians (Central) Karelians Vepsas Bashkirs
Turkmens Iranians Tajiks Pathans Uzbeks
Kyrgyz Iranians Tajiks Kazakhs
Turkic speakers grouped with background populations